CRF Err rs Corrected by the STIC Syst ms Branch

Serial Numb r: 09/926,323

CRF Processing Date: 5/20/02
Edited by: _____ VC
Verified by: _____ VC (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases wh re the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically: **ENTERED** _____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____ .

☑ Inserted mandatory headings, specifically: <220> in Seq. 2 + 4

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other: _____
_____
_____

*Examin r: Th above corrections must be communicated to th applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PCT09

```
            RAW SEQUENCE LISTING                   DATE: 05/20/2002
            PATENT APPLICATION:  US/09/926,323     TIME: 11:36:58

            Input Set : A:\PTO.DC.txt
            Output Set: N:\CRF3\05202002\I926323.raw
```

```
        3 <110> APPLICANT: Wilex Biotechnology GmbH
        5 <120> TITLE OF INVENTION: Diagnostic and therapeutic use of antibodies against
        6       the urokinase receptor
        8 <130> FILE REFERENCE: 19116pwoas
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/926,323
C--> 11 <141> CURRENT FILING DATE: 2002-04-22
       13 <150> PRIOR APPLICATION NUMBER: EP99 107 199.4
       14 <151> PRIOR FILING DATE: 1999-04-13
       16 <160> NUMBER OF SEQ ID NOS: 4
       18 <170> SOFTWARE: PatentIn Ver. 2.1
       20 <210> SEQ ID NO: 1
       21 <211> LENGTH: 354
       22 <212> TYPE: DNA
       23 <213> ORGANISM: Artificial Sequence
       25 <220> FEATURE:
       26 <223> OTHER INFORMATION: Description of Artificial Sequence: phage sequence
       28 <220> FEATURE:
       29 <221> NAME/KEY: CDS
       30 <222> LOCATION: (1)..(354)
       32 <400> SEQUENCE: 1
       33 cag gtg caa ctg cag cag tca gga cct gag ttg gtg aag cct ggg gct    48
       34 Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala
       35   1               5                  10                  15
       37 tta gtg aag ata tcc tgc aag gct tct ggt tac agt ttc aca agc tac    96
       38 Leu Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Ser Tyr
       39                20                  25                  30
       41 gat ata aat tgg gtg aag cgg agg cct gga cag gga ctt gag tgg att   144
       42 Asp Ile Asn Trp Val Lys Arg Arg Pro Gly Gln Gly Leu Glu Trp Ile
       43            35                  40                  45
       45 gga tgg att ttt cct gga gat ggt agt acc aat tac aat gag aaa ttc   192
       46 Gly Trp Ile Phe Pro Gly Asp Gly Ser Thr Asn Tyr Asn Glu Lys Phe
       47         50                  55                  60
       49 aag gac aag gcc aca ctg act gct gac aaa tcc tcc agc aca gcc tac   240
       50 Lys Asp Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr
       51 65                  70                  75                  80
       53 atg cag ctc aac agc ctg act tct gag aac tct gca gtc tat ttc tgt   288
       54 Met Gln Leu Asn Ser Leu Thr Ser Glu Asn Ser Ala Val Tyr Phe Cys
       55                85                  90                  95
       57 gca aga gat gga agt atg ggg ggg ttt gac tac tgg ggc caa ggg acc   336
       58 Ala Arg Asp Gly Ser Met Gly Gly Phe Asp Tyr Trp Gly Gln Gly Thr
       59               100                 105                 110
       61 acg gtc acc gtc tcc tca                                           354
       62 Thr Val Thr Val Ser Ser
```

RAW SEQUENCE LISTING   DATE: 05/20/2002
PATENT APPLICATION: US/09/926,323   TIME: 11:36:58

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05202002\I926323.raw

```
63          115
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 118
68 <212> TYPE: PRT
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Description of Artificial Sequence: phage sequence
74 <400> SEQUENCE: 2
75 Gln Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala
76   1               5                  10                  15
78 Leu Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Ser Phe Thr Ser Tyr
79              20                  25                  30
81 Asp Ile Asn Trp Val Lys Arg Arg Pro Gly Gln Gly Leu Glu Trp Ile
82          35                  40                  45
84 Gly Trp Ile Phe Pro Gly Asp Gly Ser Thr Asn Tyr Asn Glu Lys Phe
85      50                  55                  60
87 Lys Asp Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser Thr Ala Tyr
88  65                  70                  75                  80
90 Met Gln Leu Asn Ser Leu Thr Ser Glu Asn Ser Ala Val Tyr Phe Cys
91                  85                  90                  95
93 Ala Arg Asp Gly Ser Met Gly Gly Phe Asp Tyr Trp Gly Gln Gly Thr
94              100                 105                 110
96 Thr Val Thr Val Ser Ser
97          115
101 <210> SEQ ID NO: 3
102 <211> LENGTH: 324
103 <212> TYPE: DNA
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: Description of Artificial Sequence: phage sequence
109 <220> FEATURE:
110 <221> NAME/KEY: CDS
111 <222> LOCATION: (1)..(324)
113 <400> SEQUENCE: 3
114 gat gtt ttg atg acc caa act cca aaa ttc atg tcc aca tca gta gga    48
115 Asp Val Leu Met Thr Gln Thr Pro Lys Phe Met Ser Thr Ser Val Gly
116   1               5                  10                  15
118 gac agg gtc agc atc acc tgc aag gcc agt cag aat gtt cgt act act    96
119 Asp Arg Val Ser Ile Thr Cys Lys Ala Ser Gln Asn Val Arg Thr Thr
120              20                  25                  30
122 gta gcc tgg tat caa gag aaa cca ggg cag tct cct aaa gca ctg att   144
123 Val Ala Trp Tyr Gln Glu Lys Pro Gly Gln Ser Pro Lys Ala Leu Ile
124          35                  40                  45
126 tac ttg gca tcc aac cgg cac act gga gtc cct gat cgc ttc aca ggc   192
127 Tyr Leu Ala Ser Asn Arg His Thr Gly Val Pro Asp Arg Phe Thr Gly
128      50                  55                  60
130 agt gga tct gga aca gat ttc act ctc acc att agc aat gtg caa tct   240
131 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser
132  65                  70                  75                  80
```

RAW SEQUENCE LISTING   DATE: 05/20/2002
PATENT APPLICATION: US/09/926,323   TIME: 11:36:58

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05202002\I926323.raw

```
134 gaa gac ctg gca gat tat ttc tgt ctg caa cat tgg aat tat ccg tac    288
135 Glu Asp Leu Ala Asp Tyr Phe Cys Leu Gln His Trp Asn Tyr Pro Tyr
136                 85                  90                  95
138 acg ttc gga ggg ggc acc aag ctg gaa atc aaa cgg                    324
139 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg
140             100                 105
143 <210> SEQ ID NO: 4
144 <211> LENGTH: 108
145 <212> TYPE: PRT
146 <213> ORGANISM: Artificial Sequence
148 <220> FEATURE:
149 <223> OTHER INFORMATION: Description of Artificial Sequence: phage sequence
151 <400> SEQUENCE: 4
152 Asp Val Leu Met Thr Gln Thr Pro Lys Phe Met Ser Thr Ser Val Gly
153   1               5                  10                  15
155 Asp Arg Val Ser Ile Thr Cys Lys Ala Ser Gln Asn Val Arg Thr Thr
156                 20                  25                  30
158 Val Ala Trp Tyr Gln Glu Lys Pro Gly Gln Ser Pro Lys Ala Leu Ile
159             35                  40                  45
161 Tyr Leu Ala Ser Asn Arg His Thr Gly Val Pro Asp Arg Phe Thr Gly
162       50                  55                  60
164 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser
165  65                  70                  75                  80
167 Glu Asp Leu Ala Asp Tyr Phe Cys Leu Gln His Trp Asn Tyr Pro Tyr
168                 85                  90                  95
170 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg
171             100                 105
```

VERIFICATION SUMMARY   DATE: 05/20/2002
PATENT APPLICATION: US/09/926,323   TIME: 11:36:59

Input Set : A:\PTO.DC.txt
Output Set: N:\CRF3\05202002\I926323.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date